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RAW SEQUENCE LISTING DATE: 02/19/2002 PATENT APPLICATION: US/10/058,580 TIME: 11:10:12

Input Set : A:\98-76D1.SEQ.txt

Output Set: N:\CRF3\02192002\J058580.raw

4 5 7 9 11 11 11	<pre><110> APPLICANT: Sheppard, Paul O. Novak, Julia E. Raymond, Fenella <120> TITLE OF INVENTION: Tumor Marker Zsig62 <130> FILE REFERENCE: 98-76 <140> CURRENT APPLICATION NUMBER: US/10/058,580 <141> CURRENT FILING DATE: 2002-01-28 <160> NUMBER OF SEQ ID NOS: 8 <170> SOFTWARE: FastSEQ for Windows Version 3.0</pre>												Dorrec	Compr etter di			
	<210><211>	_	_														
	<211>				34												
	<213>				OMO	san:	iens										
	<220>				·	Dup.	LUIID										
	<221>				CDS												
	<222>					(3	316)										
24	<400>	SEÇ	UEN	CE:	L												
25	caggi	tcat	gt (catto	ccaga		_	_		-						g gga	52
26 27						Met 1	Cys	s Cys	s Tr	Pro 5	s Se:	r Pro	Tr	va.	l Glr 10	n Gly D	
29	agc (cct	ggc	att	tgg	cat	ttg	tgg	qca	gtg	ttg	gcq	tgc	cac			100
30	Ser I																
31				15				_	20					25		-	
33	cac a	_	_	_		-			_	-		-				-	148
34	His S	Ser		Ser	Arg	Gln	Gly		Leu	Arg	His	Arg		Gly	Gly	Ala	
35			30					35					40				
37	ctg						_		-	_	_					_	196
38 39	Leu I	ero 45	ser	Thr	Pro	GIY	Cys 50	Tnr	мет	Thr	ser	Thr 55	Leu	GIY	GIn	Arg	
41	ccc	_	ttσ	caa	aac	tac		gac	atc	ato	atc		CCC	gag	aaa	cat	244
42	Pro I																244
43	60				1	65					70			014	011	75	
45	tta t	tct	ttg	att	gtc	ttg	agt	gct	gca	tca	gct	aag	aca	aaa	acc	aca	292
46	Leu S																
47					80					85					90		
49							_		tgat	gagg	gat t	igtgo	caatt	it c	cggac	ccatc	346
50	Glu S	Ser	Glu	_	Lys	Lys	Thr	Ser									
51				95													406
53							-									acgtg	406
54 55																ttgct agact	466 526
56			-				-						-			etgact	526 586
57																agatg	646
- '	-) ~ 9		. , ~ (, , , ,		, , ,		, , , , , ,	, , , , ,		, - , - ;	,	יצנננ		340

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```
58
                                                                         706
    ataqtqqqtq qqtqqqqaqc tqqcqaqqqt qccccaqqca qaqqcaccqt qtqtqcaa
59 aggeetgeag gtggagaagg geetgggaet ettggagaat ggeaggaagt ttggtgtgee
                                                                         766
60 tgtagtctat gagccaggct cagggcagca aaggtctgtc ctgcaggtgt tgtqatqaqc
                                                                         826
61 tgtaccactt agtgggcacc atcaagatga acagagagta acacggtggc actgagaact
                                                                         886
62 tgagaacage teactetaga atgaactgtg teetecaaag tgtgeagage caatacetag
                                                                         946
63 gggtccccaa ggtgactgag cacgggcaca gatccagcag caaatccccc caqtccaaqa
                                                                        1006
   getgttettt ceattetetg ttettteeat tetetgttet tetggteett etgettatgg
                                                                        1066
65 caaggtgaaa gtcacaggtg gaattgteee tateacetet eecacaeeet gateteettt
                                                                        1126
66 tacaacaaag agcaagcatc ctctacaaca aagcctttgg ttggtgtcag tgcctqgctg
                                                                        1186
67 ggaggaagta actgttgttt ttactgtgtt taatttcact cctgccgtct gttcacggca
                                                                        1246
68 ccagtgatca ggttctctgc cagtgggagt gatagaaagt taccttttta aagtaaattt
                                                                        1306
69 cttggaacgc aaaaaacaag ccaagttaaa taaaaataca aaatatgggg ccaggcgcgg
                                                                        1366
70 tggctcgtgc ctgtcatccc agcactttgg gaggctgaga cggtggatca cctgaggtca
                                                                        1426
71
    ggagtttgag accagcctga ccaacaaggt gaagccccgt ctctactaaa aatacaaaaa
                                                                        1486
72 ttagccgggc gtggtggcag gcacctgtag tcccagctac tcgggaggct gggacagagg
                                                                        1546
73 aattgcttga accegggagg eggaggttge agtgageega gateaegeea ecaetgeaet
                                                                        1606
74 ccagcctggg tgacggagcg agatgccatt tcaaaacaaa aacaaaatat gtactggtac
                                                                        1666
75 cagtacacag taggaaggtg ggcaaaactt gggaaggggg atattcaaag gacagggttt
                                                                        1726
76 gggaaatgct ggatcaaggt cggggaagaa ggagaactga gaggctgtta taatttagag
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77 aagtgettet cagagtgggg gecageagee aggegeegtg geteatgeet gtaacettaa
                                                                        1846
78 cactttggga ggtctaggcg ggaggattgc ctgagcccag gagttcgagt ccagcttgtg
                                                                        1906
79
   caacatagtg agatgctgtc tctacaaaaa atttaaaaaat tagctggtgt cctctcagtg
                                                                        1966
80 tgtcttgtcc tctccatgtt tctaaaataa aggaagaaag gcccagcgca gtggcgtaca
                                                                        2026
81 cctatagtct cagcactttg ggaggccaag gtgggcagat cacttgaggt caggagttcg
                                                                        2086
82 agaccageet ggetaacatg geaaaacett gtttetactg gaaatacaaa aattagetag
                                                                        2146
83 gcgtggtggt gcacgcctgt aatcccagct acttgggagg ctgagggagg agaaccgctt
                                                                        2206
84 gageetggga ggeagagget geagtgagee aagateaeae aetgeaetee ageetgggtg
                                                                        2266
85 acagagegag actecatete aaataaataa ataaataaat aaaataaat acataaatae
                                                                        2326
86 ataaaata
                                                                        2334
88 <210> SEO ID NO: 2
89 <211> LENGTH: 99
90 <212> TYPE: PRT
91 <213> ORGANISM: Homo sapiens
93 <400> SEQUENCE: 2
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95
                     5
                                        10
96
   His Leu Trp Ala Val Leu Ala Cys His Leu Gly His Ser Ser Arq
97
                20
                                    25
                                                        30
98
   Gln Gly Ile Leu Arg His Arg Pro Gly Gly Ala Leu Pro Ser Thr Pro
99
            35
                                40
100
    Gly Cys Thr Met Thr Ser Thr Leu Gly Gln Arg Pro Leu Leu Gln Gly
101
         50
                             55
                                                 60
102 Cys Glu Asp Ile Met Val Gln Pro Glu Gly Asp Leu Ser Leu Ile Val
103
                         70
                                             75
104
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105
106 Lys Thr Ser
108 <210> SEQ ID NO: 3
109 <211> LENGTH: 297
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Input Set : A:\98-76D1.SEQ.txt

Output Set: N:\CRF3\02192002\J058580.raw

		Output Set. N:/CRF5/02192002/0036360.1aw								
	110									
		<212> TYPE: DNA (
C>		<pre><213> ORGANISM: Arificial Sequence</pre>								
		<pre><220> FEATURE:</pre>								
		<223> OTHER INFORMATION: This degenerate sequence encodes the amino a	cid							
	115									
		<221> NAME/KEY: variation								
		<222> LOCATION: (1)(297)								
		<223> OTHER INFORMATION: N is any nucleotide.								
		<400> SEQUENCE: 3								
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		gtnytngcnt gycayytngg ncaywsnwsn wsnmgncarg gnathytnmg ncaymgnccn	120							
		ggnggngcny tnccnwsnac nccnggntgy acnatgacnw snacnytngg ncarmgnccn	180							
		ytnytncarg gntgygarga yathatggtn carcengarg gngayytnws nytnathgtn	240							
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		<211> LENGTH: 23								
		<212> TYPE: DNA								
		<213> ORGANISM: Artificial Sequence								
		<220> FEATURE:								
		<223> OTHER INFORMATION: PCR primer								
		<400> SEQUENCE: 4								
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		<211> LENGTH: 24								
	141	<212> TYPE: DNA								
	142	<213> ORGANISM: Artificial Sequence								
	144	<220> FEATURE:								
		<223> OTHER INFORMATION: PCR primer								
	147	<400> SEQUENCE: 5								
	148	ggcatttgtg ggcagtgttg gggc	24							
	150	<210> SEQ ID NO: 6								
	151	<211> LENGTH: 18								
		<212> TYPE: DNA								
	153	<213> ORGANISM: Artificial Sequence								
		<220> FEATURE:								
		<223> OTHER INFORMATION: PCR primer								
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	161	<210> SEQ ID NO: 7								
	162	<211> LENGTH: 18								
		<212> TYPE: DNA								
		<213> ORGANISM: Artificial Sequence								
		<220> FEATURE:								
	167	<223> OTHER INFORMATION: PCR primer								
	169	<400> SEQUENCE: 7								
	170	tececaceca eccaetat	18							
	177	<210 SEC ID NO. 8								

172 <210> SEQ ID NO: 8 173 <211> LENGTH: 16 174 <212> TYPE: PRT

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Output Set: N:\CRF3\02192002\J058580.raw

175 <213> ORGANISM: Artificial Sequence

177 <220> FEATURE:

178 <223> OTHER INFORMATION: Peptide linker

180 <400> SEQUENCE: 8

181 Gly Gly Ser Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser

182 1

VERIFICATION SUMMARY

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Input Set : A:\98-76D1.SEQ.txt

Output Set: N:\CRF3\02192002\J058580.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:111 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3

L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

 $L\!:\!126~M\!:\!341~W\!:$ (46) "n" or "Xaa" used, for SEQ ID#:3